STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

| Application Serial Number: | 10/528, 260 A |
|---------------------------------------|---------------|
| Source: | IFWP |
| Date Processed by STIC: | 08/14/2006 |
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THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 4.4.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

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DATE: 08/14/2006

PATENT APPLICATION: US/10/528,260A

TIME: 13:29:48

Input Set : E:\P1983R1.txt

Output Set: N:\CRF4\08142006\J528260A.raw

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3 <110> APPLICANT: HENRY CHIU
              HILARY CLARK
              KATHRYN DENNIS
      6
              SHERMAN FONG
              JILL R. SCHOENFELD
              WILLIAM I. WOOD
      q
              THOMAS D. WU
     11 <120> TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATING IMMUNE
DISORDERS
     13 <130> FILE REFERENCE: P1983R1
                                                                  Does Not Comply
     15 <140> CURRENT APPLICATION NUMBER: US 10/528,260A
                                                                  Corrected Diskette Needed
     16 <141> CURRENT FILING DATE: 2005-03-16
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     18 <150> PRIOR APPLICATION NUMBER: US 60/411,392
     19 <151> PRIOR FILING DATE: 2002-09-16
     21 <150> PRIOR APPLICATION NUMBER: US03/029097
     22 <151> PRIOR FILING DATE: 2003-09-15
                                               30-) et missing sequence
described on page-2
are Intentionally skipped
they must be included
in 61607 response
     24 <160> NUMBER OF SEQ ID NOS ( 506
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E--> 42975 <210> SEQ ID NO: 429

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/14/2006 PATENT APPLICATION: US/10/528,260A

TIME: 13:29:51

Input Set : E:\P1983R1.txt

Output Set: N:\CRF4\08142006\J528260A.raw

Skipped Sequences (NEW RULES): Sequence(s) _missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

<400> sequence id number

-> See p-4 fr sample

the lbore sequence are missing

Seq#:176,224,226,234,238,246,248,252,330,332,338,342,386,404,412,416,418,424

Seq#:426,428,430,480,492,494

DATE: 08/14/2006

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/528,260A TIME: 13:29:51

Input Set : E:\P1983R1.txt

Output Set: N:\CRF4\08142006\J528260A.raw

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L:22238 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:213 after pos.:1350
L:23561 M:214 E: (33) Seq.# missing, SEQ ID NO:224
L:23579 M:214 E: (33) Seq.# missing, SEQ ID NO:226
L:23599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:100
M:341 Repeated in SeqNo=227
L:23961 M:214 E: (33) Seq.# missing, SEQ ID NO:234
L:24152 M:214 E: (33) Seq.# missing, SEQ ID NO:238
L:24194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:239 after pos.:400
M:341 Repeated in SeqNo=239
L:24938 M:214 E: (33) Seq.# missing, SEQ ID NO:246
L:25002 M:214 E: (33) Seq.# missing, SEQ ID NO:248
L:25335 M:214 E: (33) Seq.# missing, SEQ ID NO:252
L:27797 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:281 after pos.:2500
L\colon\!33378 M\colon\!341 W\colon (46) "n" or "Xaa" used, for SEQ ID\!\#\colon\!321 after pos.:2700
L:33599 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:322 after pos.:900
L:34738 M:214 E: (33) Seq.# missing, SEQ ID NO:330
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L:34773 M:214 E: (33) Seq.# missing, SEQ ID NO:332
L:35071 M:214 E: (33) Seq.# missing, SEQ ID NO:338
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L:41386 M:214 E: (33) Seq.# missing, SEQ ID NO:404
L:42010 M:214 E: (33) Seq.# missing, SEQ ID NO:412
L:42355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:415 after pos.:0
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L:42422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:417 after pos.:750
L:42426 M:214 E: (33) Seq.# missing, SEQ ID NO:418
L:42861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:423 after pos.:0
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L:42897 M:214 E: (33) Seq.# missing, SEQ ID NO:424
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L:49277 M:214 E: (33) Seq.# missing, SEQ ID NO:480
L:50541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:491 after pos.:1950
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L:50859 M:214 E: (33) Seq.# missing, SEQ ID NO:494
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L:51187 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:496 after pos.:0
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(end of Sequence 175)

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Ser Asn Where is Sequence 176?

<210> 177

<211> 2533

<212> DNA

<213> Homo sapiens

<400> 177

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(Sample of numerous missing Sequence listing file). Throughout Submitted Sequence listing file).